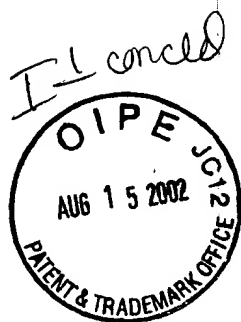


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CECH *et al.*  
Application No.: 08/974,584  
Page 3

amino acid selected from Ala, Leu, Ile, Val, Pro, Phe, Trp, and Met; and p is a polar amino acid selected from Gly, Ser, Thr, Tyr, Cys, Asn and Gln.

121. (Amended) The polynucleotide of claim 119, where structure a) further comprises Arg-Lys-X<sub>2</sub>-Trp-X<sub>2</sub>-Leu (SEQ ID NO:477).

*I2*

122. (Amended) The polynucleotide of claim 119, where structure b) comprises h-Arg-h-X-Pro-Lys (SEQ ID NO:473), wherein h is a hydrophobic amino acid selected from Ala, Leu, Ile, Val, Pro, Phe, Trp, and Met.

123. (Amended) The polynucleotide of claim 119, where structure c) comprises Arg-X-Ile-Pro-Lys (SEQ ID NO:478).

124. (Amended) The polynucleotide of claim 119, where structure e) comprises Gly-Ile-Pro-Gln-Gly-Ser (SEQ ID NO:370).

125. (Amended) The polynucleotide of claim 119, where structure f) comprises Leu-Leu-Leu-Arg-Leu-X-Asp-Asp-Phe-Leu (SEQ ID NO:479).

#### REMARKS

Claims 119-127 are pending in this application. Claims 119-125 have been amended. The amendments to claims 119-125 insert the assigned identifiers for sequences designated in these claims.

In addition, amended use of subscripts to replace superscripts to designate certain amino acid residues of the motifs recited in claim 119 brings their usage into conformity with their exact format in the Specification, *e.g.*, page 26, lines 1-2 and page 42, lines 14 and 19.